FIGURE 1

GGCGTCCGCGCACACCTCCCCGCGCCGCCGCCGCCACCGCCCGCACTCCGCCGCCTCTGCCCGCAACCGCT
GAGCCATCCATGGGGGTCGCGGGCCGCAACCGTCCCGGGGCGGCCTGGGCGGTGCTGCTGCTGCTGCTGCT
GCTGCCGCCACTGCTGCTGCTGGCGGGGGCCGTCCCGCCGGGTCGGGGCCGTGCCGCGGGGCCGCAGGAGG
ATGTAGATGAGTGTGCCCAAGGGCTAGATGACTGCCATGCCGACGCCCTGTGTCAGAACACACCCACCTCC
TACAAGTGCTCCTGCAAGCCTGGCTACCAAGGGGAAGGCAGGCAGTGTGAGGACATCGATGAATGTGGAAA
TGAGCTCAATGGAGGCTGTGTCCATGACTGTTTGAATATTCCAGGCAATTATCGTTGCACTTGTTTTGATG
GCTTCATGTTGGCTCATGACGGTCATAATTGTCTTGATGTGGACGAGTGCCTGGAGAACAATGGCGGCTGC
CAGCATACCTGTGTCAACGTCATGGGGAGCTATGAGTGCTGCTGCAAGGAGGGGTTTTTCCTGAGTGACAA
TCAGCACACCTGCATTCACCGCTCGGAAGAGGGCCTGAGCTGCATGAATAAGGATCACGGCTGTAGTCACA
TCTGCAAGGAGGCCCCAAGGGGCAGCGTCGCCTGTGAGTGCAGGCCTGGTTTTGAGCTGGCCAAGAACCAG
AGAGACTGCATCTTGACCTGTAACCATGGGAACGGTGGGTGCCAGCACTCCTGTGACGATACAGCCGATGG
CCCAGAGTGCAGCTGCCATCCACAGTACAAGATGCACACAGATGGGAGGAGCTGCCTTGAGCGAGAGGACA
CTGTCCTGGAGGTGACAGAGAGCAACACCACATCAGTGGTGGATGGGGATAAACGGGTGAAACGGCGGCTG
CTCATGGAAACGTGTGCTGTCAACAATGGAGGCTGTGACCGCACCTGTAAGGATACTTCGACAGGTGTCCA
CTGCAGTTGTCCTGTTGGATTCACTCTCCAGTTGGATGGGAAGACATGTAAAGATATTGATGAGTGCCAGA
CCCGCAATGGAGGTTGTGATCATTTCTGCAAAAACATCGTGGGCAGTTTTGACTGCGGCTGCAAGAAAGGA
TTTAAATTATTAACAGATGAGAAGTCTTGCCAAGATGTGGATGAGTGCTCTTTGGATAGGACCTGTGACCA
CAGCTGCATCAACCACCCTGGCACATTTGCTTGTGCTTGCAACCGAGGGTACACCCTGTATGGCTTCACCC
ACTGTGGAGACACCAATGAGTGCAGCATCAACAACGGAGGCTGTCAGCAGGTCTGTGTGAACACAGTGGGC
AGCTATGAATGCCAGTGCCACCCTGGGTACAAGCTCCACTGGAATAAAAAAGACTGTGTGTGGAAGTGAAGGG
GCTCCTGCCCACAAGTGTGTCACCCCGTGTGTCCCTGCACTGCGGTAAGAGTGGTGGAGGAGACGGGTGCT
TCCTCAGATGTCACTCTGGCATTCACCTCTCTTCAGATGTCACCACCATCAGGACAAGTGTAACCTTTAAG
CTAAATGAAGGCAAGTGTAGTTTGAAAAATGCTGAGCTGTTTCCCGAGGGTCTGCGACCAGCACTACCAGA
GAAGCACAGCTCAGTAAAAGAGAGCTTCCGCTACGTAAACCTTACATGCAGCTCTGGCAAGCAAGTCCCAG
GAGCCCCTGGCCGACCAAGCACCCCTAAGGAAATGTTTATCACTGTTGAGTTTGAGCTTGAAACTAACCAA
AAGGAGGTGACAGCTTCTTGTGACCTGAGCTGCATCGTAAAGCGAACCGAGAAGCGGCTCCGTAAAGCCAT
CCGCACGCTCAGAAAGGCCGTCCACAGGGAGCAGTTTCACCTCCAGCTCTCAGGCATGAACCTCGACGTGG
CTAAAAAGCCTCCCAGAACATCTGAACGCCAGGCAGAGTCCTGTGGAGTGGGCCAGGGTCATGCAGAAAAC
CAATGTGTCAGTTGCAGGGCTGGGACCTATTATGATGGAGCACGAGAACGCTGCATTTTATGTCCAAATGG
AACCTTCCAAAATGAGGAAGGACAAATGACTTGTGAACCATGCCCAAGACCAGGAAATTCTGGGGCCCTGA
AGACCCCAGAAGCTTGGAATATGTCTGAATGTGGAGGTCTGTGTCAACCTGGTGAATATTCTGCAGATGGC
TTTGCACCTTGCCAGCTCTGTGCCCTGGGCACGTTCCAGCCTGAAGCTGGTCGAACTTCCTGCTTCCCCTG
TGGAGGAGGCCTTGCCACCAAACATCAGGGAGCTACTTCCTTTCAGGACTGTGAAACCAGAGTTCAATGTT
CACCTGGACATTTCTACAACACCACCACTCACCGATGTATTCGTTGCCCAGTGGGAACATACCAGCCTGAA
TTTGGAAAAAATAATTGTGTTTCTTGCCCAGGAAATACTACGACTGACTTTGATGGCTCCACAAACATAAC
CCAGTGTAAAAACAGAAGATGTGGAGGGGAGCTGGGAGATTTCACTGGGTACATTGAATCCCCAAACTACC
CAGGCAATTACCCAGCCAACACCGAGTGTACGTGGACCATCAACCCACCCCCCAAGCGCCGCATCCTGATC
GTGGTCCCTGAGATCTTCCTGCCCATAGAGGACGACTGTGGGGACTATCTGGTGATGCGGAAAACCTCTTC
ATCCAATTCTGTGACAACATATGAAACCTGCCAGACCTACGAACGCCCCATCGCCTTCACCTCCAGGTCAA
AGAAGCTGTGGATTCAGTTCAAGTCCAATGAAGGGAACAGCGCTAGAGGGTTCCAGGTCCCATACGTGACA
TATGATGAGGACTACCAGGAACTCATTGAAGACATAGTTCGAGATGGCAGGCTCTATGCATCTGAGAACCA
TCAGGAAATACTTAAGGATAAGAAACTTATCAAGGCTCTGTTTGATGTCCTGGCCCATCCCCAGAACTATT
TCAAGTACACAGCCCAGGAGTCCCGAGAGATGTTTCCAAGATCGTTCATCCGATTGCTACGTTCCAAAGTG
TCCAGGTTTTTGAGACCTTACAAATGACTCAGCCCACGTGCCACTCAATACAAATGTTCTGCTATAGGGTT
GGTGGGACAGAGCTGTCTTCCTTCTGCATGTCAGCACAGTCGGGTATTGCTGCCTCCCGTATCAGTGACTC
ATTAGAGTTCAATTTTTATAGATAATACAGATATTTTGGTAAATTGAACTTGGTTTTTCTTTCCCAGCATC
GTGGATGTAGACTGAGAATGGCTTTGAGTGGCATCAGCTTCTCACTGCTGTGGGCGGATGTCTTGGATAGA
TCACGGGCTGGCTGAGCTGGACTTTGGTCAGCCTAGGTGAGACTCACCTGTCCTTCTGGGGTCTTACTCCT
CCTCAAGGAGTCTGTAGTGGAAAGGAGGCCACAGAATAAGCTGCTTATTCTGAAACTTCAGCTTCCTCTAG
CCCGGCCCTCTCTAAGGGAGCCCTCTGCACTCGTGTGCAGGCTCTGACCAGGCAGAACAGGCAAGAGGGGA
GGGAAGGAGACCCCTGCAGGCTCCCTCCACCCACCTTGAGACCTGGGAGGACTCAGTTTCTCCACAGCCTT
CTCCAGCCTGTGTGATACAAGTTTGATCCCAGGAACTTGAGTTCTAAGCAGTGCTCGTGAAAAAAAAAAGC
AGAAAGAATTAGAAATAAATAAAAACTAAGCACTTCTGGAGACAT

FIGURE 2

MGVAGRNRPGAAWAVLLLLLLLPPLLLLAGAVPPGRGRAAGPQEDVDECAQGLDDCHADALCQNTPTSYKC
SCKPGYQGEGRQCEDIDECGNELNGGCVHDCLNIPGNYRCTCFDGFMLAHDGHNCLDVDECLENNGGCQHT
CVNVMGSYECCCKEGFFLSDNQHTCIHRSEEGLSCMNKDHGCSHICKEAPRGSVACECRPGFELAKNQRDC
ILTCNHGNGGCQHSCDDTADGPECSCHPQYKMHTDGRSCLEREDTVLEVTESNTTSVVDGDKRVKRRLLME
TCAVNNGGCDRTCKDTSTGVHCSCPVGFTLQLDGKTCKDIDECQTRNGGCDHFCKNIVGSFDCGCKKGFKL
LTDEKSCQDVDECSLDRTCDHSCINHPGTFACACNRGYTLYGFTHCGDTNECSINNGGCQQVCVNTVGSYE
CQCHPGYKLHWNKKDCVEVKGLLPTSVSPRVSLHCGKSGGGDGCFLRCHSGIHLSSDVTTIRTSVTFKLNE
GKCSLKNAELFPEGLRPALPEKHSSVKESFRYVNLTCSSGKQVPGAPGRPSTPKEMFITVEFELETNQKEV
TASCDLSCIVKRTEKRLRKAIRTLRKAVHREQFHLQLSGMNLDVAKKPPRTSERQAESCGVGQGHAENQCV
SCRAGTYYDGARERCILCPNGTFQNEEGQMTCEPCPRPGNSGALKTPEAWNMSECGGLCQPGEYSADGFAP
CQLCALGTFQPEAGRTSCFPCGGGLATKHQGATSFQDCETRVQCSPGHFYNTTTHRCIRCPVGTYQPEFGK
NNCVSCPGNTTTDFDGSTNITQCKNRRCGGELGDFTGYIESPNYPGNYPANTECTWTINPPPKRRILIVVP
EIFLPIEDDCGDYLVMRKTSSSNSVTTYETCQTYERPIAFTSRSKKLWIQFKSNEGNSARGFQVPYVTYDE
DYQELIEDIVRDGRLYASENHQEILKDKKLIKALFDVLAHPQNYFKYTAQESREMFPRSFIRLLRSKVSRF
LRPYK

FIGURE 3

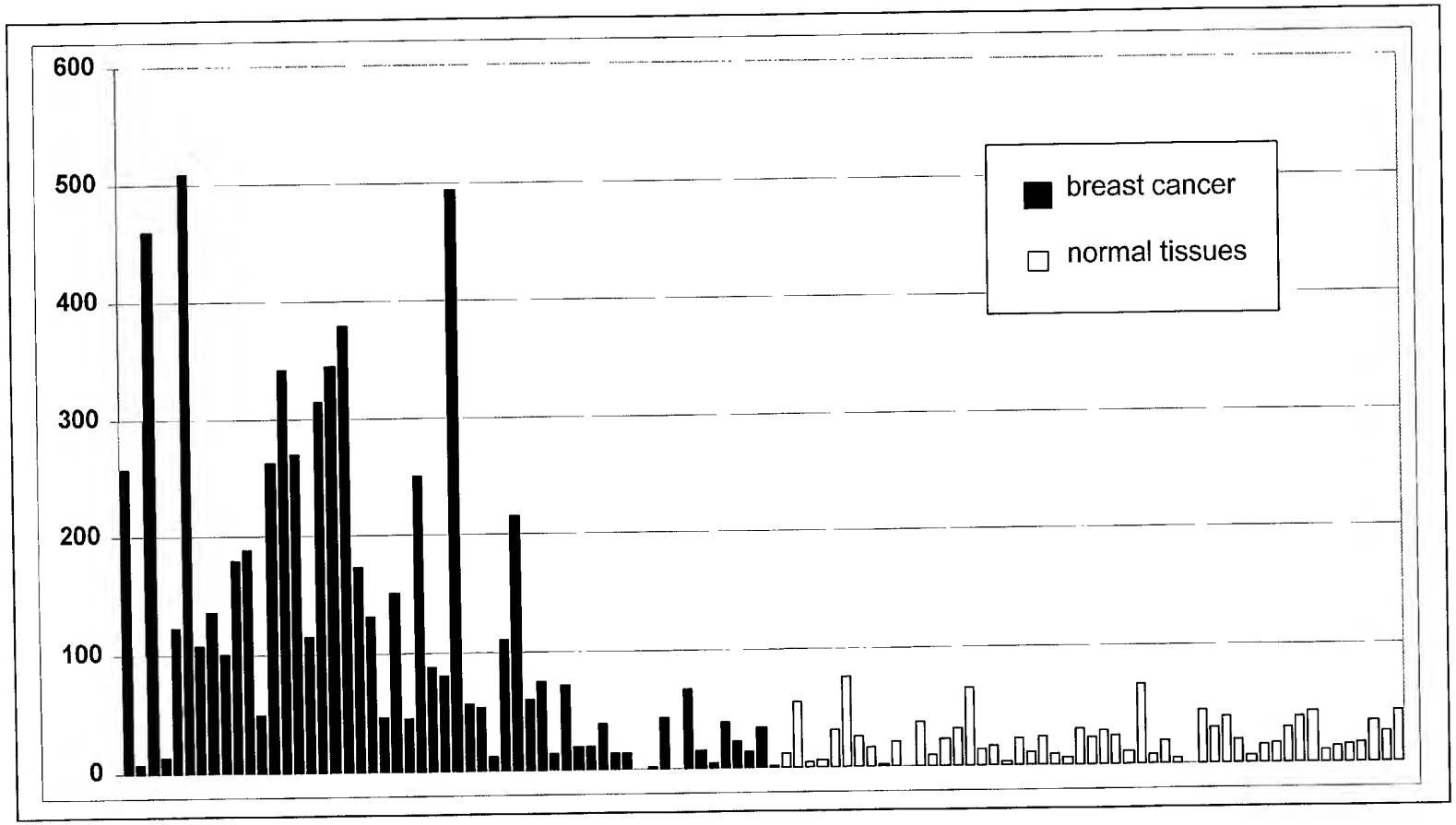

FIGURE 4A

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BCO2_human    MGVAGRNRPGAAWAVLLLLLLLPPLLLLAGAVPPGRGRAAGPQEDVDECAQGLDDCHADA
BCO2_mouse    MGVAGCGRPREARALLLLLLLLPP--LLAAAVPPDRGLTNGPSEDVDECAQGLDDCHADA
              ***** .**  * *:*********  ***.****.** :  **.*****************

BCO2_human    LCQNTPTSYKCSCKPGYQGEGRQCEDIDECGNELNGGCVHDCLNIPGNYRCTCFDGFMLA
BCO2_mouse    LCQNTPTSYKCSCKPGYQGEGRQCEDMDECDNTLNGGCVHDCLNIPGNYRCTCFDGFMLA
              **************************:***.* ***************************

BCO2_human    HDGHNCLDVDECLENNGGCQHTCVNVMGSYECCCKEGFFLSDNQHTCIHRSEEGLSCMNK
BCO2_mouse    HDGHNCLDMDECLENNGGCQHICTNVIGSYECRCKEGFFLSDNQHTCIHRSEEGLSCMNK
              ********:************ *.**:***** ***************************

BCO2_human    DHGCSHICKEAPRGSVACECRPGFELAKNQRDCILTCNHGNGGCQHSCDDTADGPECSCH
BCO2_mouse    DHGCGHICKEAPRGSVACECRPGFELAKNQKDCILTCNHGNGGCQHSCEDTAEGPECSCH
              ****.*************************:******************:***:******

BCO2_human    PQYKMHTDGRSCLEREDTVLEVTESNTTSVVDGDKRVKRRLLMETCAVNNGGCDRTCKDT
BCO2_mouse    PRYRLHADGRSCLEQEGTVLEGTESNATSVADGDKRVKRRLLMETCAVNNGGCDRTCKDT
              *:*::*:*******:*.**** ****:***.*****************************

BCO2_human    STGVHCSCPVGFTLQLDGKTCKDIDECQTRNGGCDHFCKNIVGSFDCGCKKGFKLLTDEK
BCO2_mouse    STGVHCSCPTGFTLQVDGKTCKDIDECQTRNGGCNHFCKNTVGSFDCSCKKGFKLLTDEK
              *********.*****:******************:***** ******.************

BCO2_human    SCQDVDECSLDRTCDHSCINHPGTFACACNRGYTLYGFTHCGDTNECSINNGGCQQVCVN
BCO2_mouse    SCQDVDECSLERTCDHSCINHPGTFICACNPGYTLYSFTHCGDTNECSVNNGGCQQVCIN
              **********:************** **** *****.***********:*********:*

BCO2_human    TVGSYECQCHPGYKLHWNKKDCVEVKGLLPTSVSPRVSLHCGKSGGGDGCFLRCHSGIHL
BCO2_mouse    TVGSYECQCHPGFKLHWNKKDCVEVKGFPPTSMTPRVSLHCGKSGGGDRCFLRCRSGIHL
              ************:**************: ***::************** *****:*****

BCO2_human    SSDVTTIRTSVTFKLNEGKCSLKNAELFPEGLRPALPEKHSSVKESFRYVNLTCSSGKQV
BCO2_mouse    SSDVVTVRTSVTFKLNEGKCSLQKAKLSPEGLRPALPERHSSVKESFQYANLTCSPGKQV
              ****.*:***************::*:* **********:********:*.*****.****

BCO2_human    PGAPGRPSTPKEMFITVEFELETNQKEVTASCDLSCIVKRTEKRLRKAIRTLRKAVHREQ
BCO2_mouse    PGALGRLNAPKEMFITVEFERETYEKEVTASCNLSCVVKRTEKRLRKALRTLKRAAHREQ
              *** ** .:*********** ** :*******:***:***********:***::*.****

BCO2_human    FHLQLSGMNLDVAKKPPRTSERQAESCGVGQGHAENQCVSCRAGTYYDGARERCILCPNG
BCO2_mouse    FHLQLSGMDLDMAKTPSRVSGQHEETCGVGQGHEESQCVSCRAGTYYDGSQERCILCPNG
              ********:**:**.*.*.* :: *:******* *.*************::*********

BCO2_human    TFQNEEGQMTCEPCPRPGNSGALKTPEAWNMSECGGLCQPGEYSADGFAPCQLCALGTFQ
BCO2_mouse    TFQNEEGQVTCEPCPRPENLGSLKISEAWNVSDCGGLCQPGEYSANGFAPCQLCALGTFQ
              ********:******** * *:** .****:*:************:**************

BCO2_human    PEAGRTSCFPCGGGLATKHQGATSFQDCETRVQCSPGHFYNTTTHRCIRCPVGTYQPEFG
BCO2_mouse    PDVGRTSCLSCGGGLPTKHLGATSFQDCETRVQCSPGHFYNTTTHRCIRCPLGTYQPEFG
              *:.*****:.*****.*** *******************************:********

BCO2_human    KNNCVSCPGNTTTDFDGSTNITQCKNRRCGGELGDFTGYIESPNYPGNYPANTECTWTIN
BCO2_mouse    KNNCVSCPGNTTTDFDGSTNITQCKNRKCGGELGDFTGYIESPNYPGNYPANSECTWTIN
              ***************************:************************:*******

BCO2_human    PPPKRRILIVVPEIFLPIEDDCGDYLVMRKTSSSNSVTTYETCQTYERPIAFTSRSKKLW
BCO2_mouse    PPPKRRILIVVPEIFLPIEDDCGDYLVMRKTSSSNSVTTYETCQTYERPIAFTSRSKKLW
              ************************************************************
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BCO2_human      IQFKSNEGNSARGFQVPYVTYDEDYQELIEDIVRDGRLYASENHQEILKDKKLIKALFDV
BCO2_mouse      IQFKSNEGNSARGFQVPYVTYDEDYQELIEDIVRDGRLYASENHQEILKDKKLIKALFDV
                ************************************************************

BCO2_human      LAHPQNYFKYTAQESREMFPRSFIRLLRSKVSRFLRPYK
BCO2_mouse      LAHPQNYFKYTAQESREMFPRSFIRLLRSKVSRFLRPYK
                ***************************************
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FIGURE 4B